

Fig. 1

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMPSCDYSTSTEELTSLRIYWQKDSKWVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQXDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKWVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWRKDSKMXLAILP MGHTMKWGSLPPKCPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKWVLAILP MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEKLTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSL PPKRPCLWL PQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRI YWQKDSKWVLAILP MGHTMKWGSL PPKR PCLMLSQLLVLTGL FYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWPSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKWVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWRSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWRSL PPKR PCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNASTEELTSLRIYWQKDSKWVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKWVLAILP MGHTMKWGSL PPKR PCLWL SQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTLRPGTPLPRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWQKDQQMVLSIIS MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTLRPGTPLPRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWQKDQQMVLSIIS MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETTVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTRRQGTSPSKCPYLNFFQLLVLAGLSHFCSG--VIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMS MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTLRPGTPLPRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWQKDQQMVLSIIS Extracellular domain (ECD) Signal sequence 222 $\widehat{\Xi}$ (1) (1) (1) 1 <u>1</u> 1 <u>1</u> $\widehat{\Xi}$ (1) (1) <u>-</u> Ξ 1) (T) 3 (1) 1 (1) 200 2222 (1) SEQ:067_R2_CD28BP-16 SEQ:175_cd28a4-5star SEQ:176_cd28A4-9 SEQ:177_cd28A6-9 SEQ:180_cd28A8-6 SEQ:181_cd28B2-8 SEQ:182_cd28B4-3 SEQ:183_cd28B6-3 SEQ:184_cd28b6-6 SEQ:185_cd28b8-5star SEQ:186_cd28c11-5 SEQ:061_R2_CD28BP-10 SEQ:066_R2_CD28BP-15 SEQ:174_cd28A12-5 SEQ:178_cd28A6-1 SEQ: 179_cd28A8-4 SEQ:050_R1_Clone_118 SEQ:051_R1_Clone_126 SEQ:056_R2_CD28BP-5 SEQ:057_R2_CD28BP-6 SEQ:059_R2_CD28BP-8 SEQ:060_R2_CD28BP-9 SEQ:063_R2_CD28BP-12 SEQ:064_R2_CD28BP-13 SEQ:065_R2_CD28BP-14 SEQ:049_R1_Clone_84 SEQ:052_R2_CD28BP-1 SEQ:053_R2_CD28BP-2 SEQ:054_R2_CD28BP-3 SEQ:055_R2_CD28BP-4 SEQ:062_R2_CD28BP-11 SEQ:068_R2_CD28BP-17 SEQ:278_Human_B7-1 SEQ:048_R1_Clone_71 SEQ:058_R2_CD28BP-7

Fig. 2A

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→ Signal sequence →	MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGTTPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKWVLAILP) MGHIMKWGSLPPKRFCLWLSQLLVLIGLFYFCSGLIPRSVIRKVKEIVMLSCUXNISTEELISLKIYWQKLDSKMVLALLP) MGHTMKWGSLPPKRPCLWLSQLLVLIGLFYFCSGTIFRSVIRKVKEIVMLSCDYSTSTEELTSLK1YWQKLSKWVLALLP MCHHT BDCHDT DBCT HTVT CTTT 1 1 1 2 CT HESCC = - TSOXWWCKKYRW 3 1 SCDXNTSTDR1 3 BWRTXMOXDOOMG STTS) MGHTHAKEGI FLEKCLINDKILGOLIMI SOGISĞY IKSYKETIMI SCOLIMISTERINGELIMIKE INGENESINDEN MENDEKINDEN MENDE MENDEKINDEN MENDEKINDEN MENDEK) MGHTMKWGSLPPKRPCLWLSOLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWOKDSKWVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMEWGSLPPKRPCLWLSQLLVTLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVTJGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELJSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLV1TGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTDLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVTLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP	_	_	_	_	_	_	_		_	_			1) MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP
	Ξ	ਹ :	_ ;	J 5	j :	ਹੋ ਹੋ	<u>ל</u>	7	<u></u>	1	7	1	(1	(1)	1	7	7	(1	7	(1	(1	(1	1	1)	1	(1	1	1	7	(1	(1	[]	(1)	7	(1
	SEQ:187_cd28C6-1	SEQ: 188_cd28C7-3	SEQ: 189_cd28C8-6	SEQ:190_cd28c9-5star	SEO:191_CG26C2-4 SEO:192 CG28C2-3	SEO:193 cd28D2-9	SEQ: 194_cd28D8-9	SEQ:195_cd28D11-1	SEQ:196_cd28D12-5	SEQ:197_cd28E10-6	SEQ:198_cd28F7-2	SEQ:199_cd28F8-4	SEQ:200_cd28F10-2	SEQ:201_cd28F12-5star	SEQ:202_cd28G2-8	SEQ:203_cd28G1-5	SEQ:204_cd28G1-9	SEQ:205_cd28H4-3	SEQ:206_cd28H11-3	SEQ:207_cd28H6-6	SEQ:208_cd28E2-4	SEQ:209_cd28B4-5a	SEQ:210_cd28A2-5a	SEQ:211_cd28B4-5star	SEQ:212_cd28D5-6	SEQ:213_cd28D10-4	SEQ:214_cd28E2-5star	SEQ:215_cd28E5-2	SEQ:216_cd28E8-6	SEQ:217_cd28E9-6	SEQ:218_cd28F3-1	SEQ:219_cd28F3-5	SEQ:220_cd28F3-6	SEQ:221_cd28F11-8	SEQ:283_CD28BP_Con

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Fig. 2B

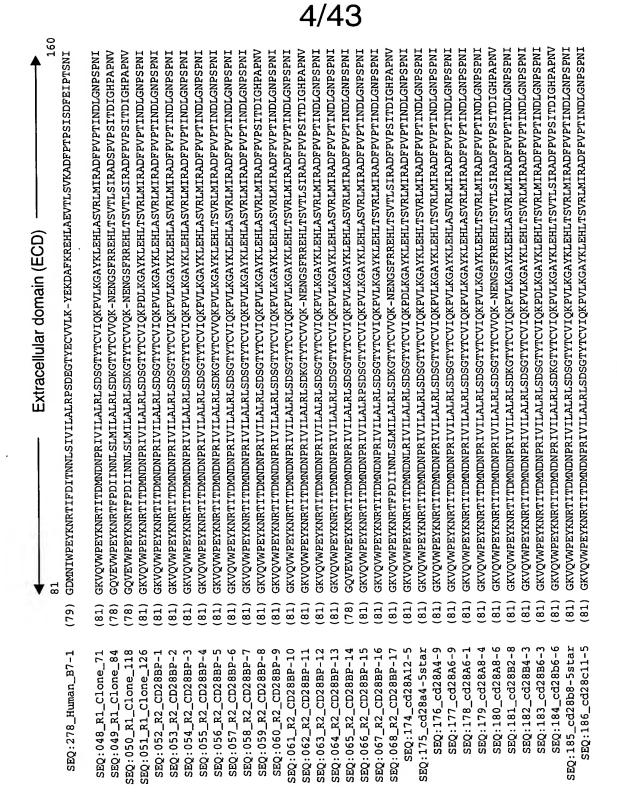


Fig. 2C

Fig. 2D

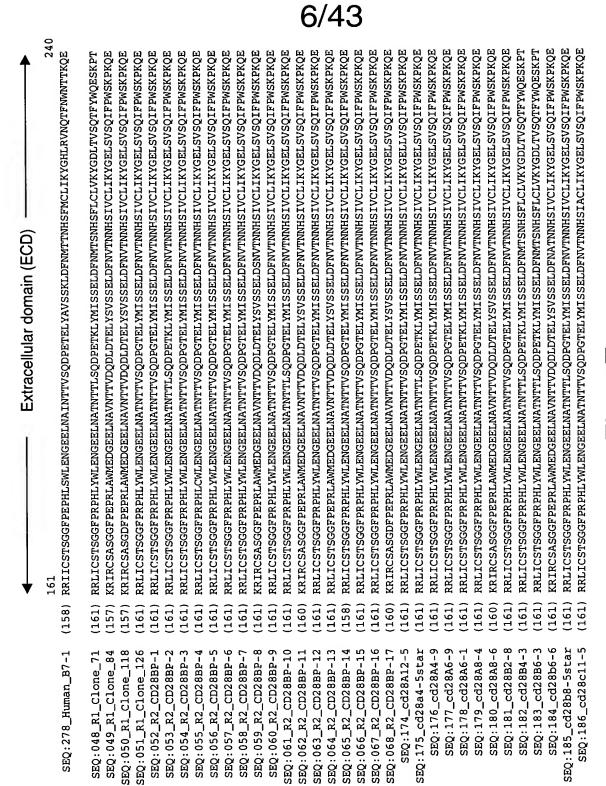


Fig. 2E

Fig. 2F

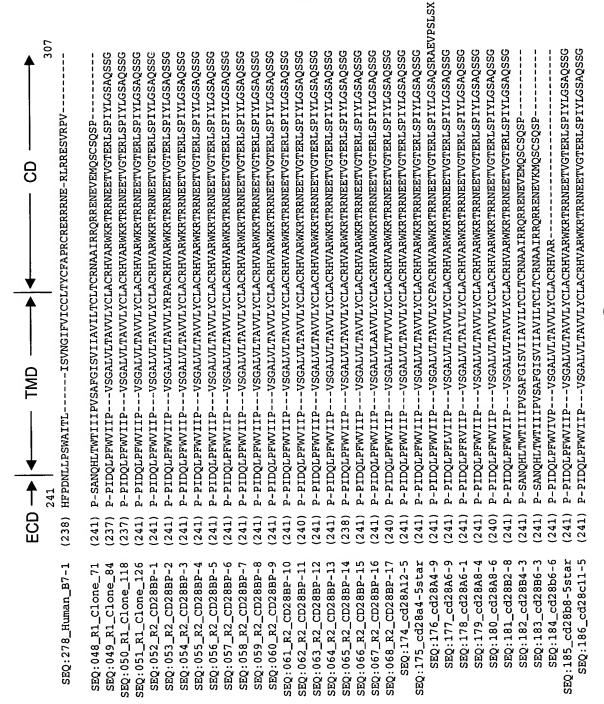


Fig. 2G

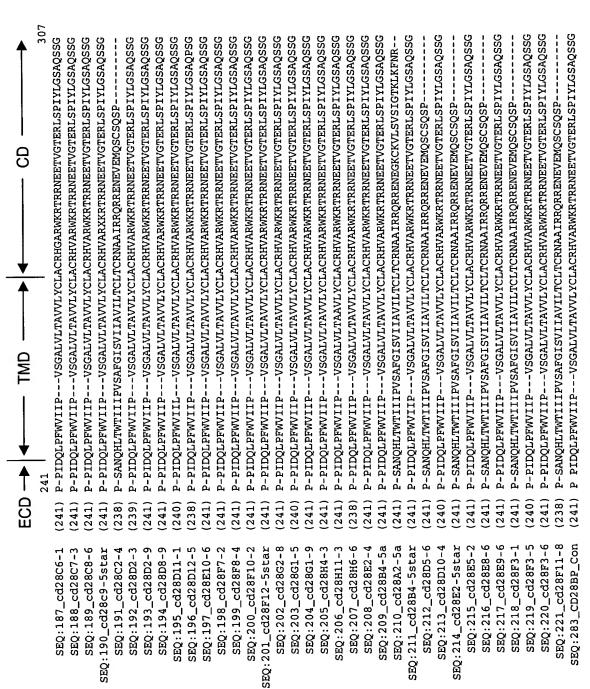


Fig. 24

(1) MGHTRRQGTSPSKCPYLNFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD

SEQ:069_R1_CTLA4BP-5 SEQ:278_Human_B7-1 SEQ: 070_R1_CTLA4BP-7 SEQ: 072_R1_CTLA4BP-13 SEQ: 073_R1_CTLA4BP-27

 Ξ <u>-</u> Ξ (1) (1)

SEQ: 071_R1_CTLA4BP-11

SEQ: 075_R2_CTLA4BP-5x2-11d SEQ: 074_R2_CTLA4BP-5x2-10c SEQ:076_R2_CTLA4BP-5X2-12F

SEQ:080_R2_CTLA4BP-5x2-7b SEQ:078_R2_CTLA4BP-5x2-3c SEQ: 079_R2_CTLA4BP-5x2-4c SEQ:077_R2_CTLA4BP-5x2-2g

SEQ:082_R2_CTLA4BP-5x3-10e SEQ:083_R2_CTLA4BP-5x3-11b SEQ:081_R2_CTLA4BP-5x2-8c

SEQ:085_R2_CTLA4BP-5x4-11d SEQ: 086_R2_CTLA4BP-5x4-12c SEQ:084_R2_CTLA4BP-5x3-6f

SEQ:088_R2_CTLA4BP-5x5-2e SEQ:087_R2_CTLA4BP-5x4-1f SEQ:089_R2_CTLA4BP-5x5-6e

SEQ:090_R2_CTLA4BP-5x6-9d SEQ:091_R2_CTLA4BP-5x8-1f SEQ: 092_R2_CTLA4BP-5x9-12c

SEQ:222_ctla5x9d10 SEQ:223_ctla5x6f6 SEQ:224_ctla5x5h12 SEQ: 225_ctla5x5c10

MGHTRRQGISPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD

MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGYTRRQGTSPSKCPYLKFFQLLVLAGLSHLCSGVIHVTNEVKEVATLSCGHNVSGEELAQTRIYWQKEKKMVLTMMYGD MSHTRROGTSPSKCPYLKFFQLLVLASLSHFCSGVIHMTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD

MSHTRRQGISPSKCPYLNFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPPKCPYLNFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD

MSHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD

MGHTRRQGTSPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGTSPSKCPYLKFFQLLVLACLSHFCSGVIHVTREVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD (1)

MSHTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIYWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLNFFRLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHMTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD $\overline{1}$ (1)(1)

MGYTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLNFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKGKKMVLTMMSGD (1) 1 (1)

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MGHTRRQGISPSKCPYLKFFQLLVLACLSHLCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGTSPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD (1) (1)

MGHTRRQGTSPSKCPYLKFFQLLVMACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIHWQKEKKMVLTMMSGD (1) (7)

MGHTRRQGISPSKCPYLKFFQLLVLAGLPHLCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGTSPSKCPYLKFFQLLVLAGLSHLCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD 1 <u>-</u>

MGHTRRQGISPSKCPYLNFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGTSPSKCPYLNFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD 1

MSHTRRQGTSPSKCPYLKFFQFLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIYWQKGKKMVLTMMSGD <u>-</u> Ξ

MGYTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTPIYWQKEKKMVLTMMSGD MGYTRRQGISPSKCPYLKFFQLLVLASLSHFCSGVIHVTKKVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD (1)(1)

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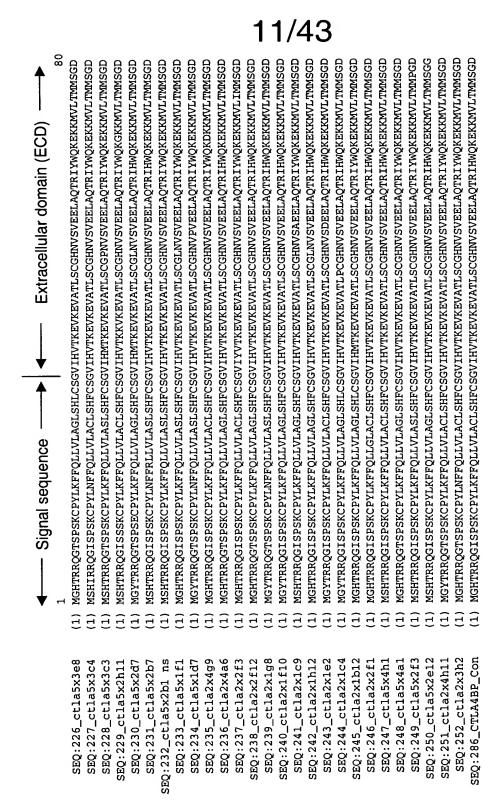


Fig. 3B

Fig. 3C

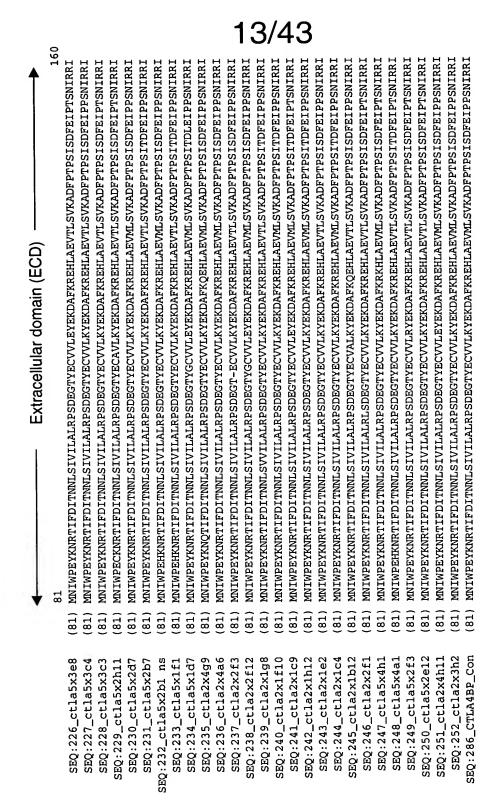


Fig. 3D

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◆ Extracellular domain (ECD)	161 ICSTSGGFPEPHLSWLENGEELNAINT	1 ICSTSGGFPEPHLSWLENGEELNAINTTVVSQDPGTELYAVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 1 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYAVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 2 ICSTSGGFPEPHLSWLENGEEINAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 3 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 3 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 3 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 4 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 5 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 6 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 7 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTTNHSFWCLIKYGHLRVNQTFNWNTPKQEHFP 8 ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTVSSKLDFNWTNH
	(161)	(161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161) (161)
	SEQ:278_Human_B7-1	SEQ: 069_R1_CTLA4BP-5 SEQ: 070_R1_CTLA4BP-7 SEQ: 071_R1_CTLA4BP-13 SEQ: 072_R1_CTLA4BP-13 SEQ: 073_R1_CTLA4BP-57 SEQ: 074_R2_CTLA4BP-5x2-10c SEQ: 075_R2_CTLA4BP-5x2-11d SEQ: 076_R2_CTLA4BP-5x2-12f SEQ: 076_R2_CTLA4BP-5x2-12f SEQ: 076_R2_CTLA4BP-5x2-12f SEQ: 078_R2_CTLA4BP-5x2-12f SEQ: 081_R2_CTLA4BP-5x2-16 SEQ: 081_R2_CTLA4BP-5x2-16 SEQ: 081_R2_CTLA4BP-5x3-11b SEQ: 081_R2_CTLA4BP-5x3-16 SEQ: 082_R2_CTLA4BP-5x3-11c SEQ: 086_R2_CTLA4BP-5x3-11c SEQ: 086_R2_CTLA4BP-5x4-12c SEQ: 086_R2_CTLA4BP-5x4-12c SEQ: 086_R2_CTLA4BP-5x4-11c SEQ: 086_R2_CTLA4BP-5x4-11c SEQ: 090_R2_CTLA4BP-5x6-9d SEQ: 092_R2_CTLA4BP-5x9-12c SEQ: 222_CTLa5x9610 SEQ: 222_CTLa5x9510

Fig. 3E

Fig. 3F

(241) DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVRPV

SEQ:278_Human_B7-1

DNLL PSWAITLISVNGIFVICCLTHCFA PRCRERRNERLRRESARPV DNLL PSWAITLISANGIFVICCLTYRFAPRCRERRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNETLRRESVRPV ONLL PSWAITLISVNGIFVICCLTYCFAPRCRERR-NETLRRESVRPV DNPL PSWAITLISANGIFVICCLTYCFA PRCRERRNETLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYRFAPRCRERKSNERLRRESVRPV ONLL PSWAITLISANGIFVICCLTYCFAPRCRERRNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERRRNERLRRESVHPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRSNERLRRESVRPV DNLLPSWAITLISANGIFVICCLAYCFAPGCRERKSNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYRFAPRCRERKSNETLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNETLRRESVRPV DNLL PSWAITLISANGIFVICCLTHCFAPRCRERKRNERLRRESVRPV DNLL PSWAITL I SVNGIFVICCL TYCFA PRCRERRRNERLRRESVR PV DNLL PSWAITLISVNGIFVICCL TYCFAPRCRERKSNERLRRESVRPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERKSNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVRPV DNLL PSWAITLISANGIFVICCLTYCFAPRCRERKSNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERKSNETLRRESVRPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVHPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERKSNERLRRESVHPV DNLL PSWAITLISVNGIFVICCLTYRFAPRCRERKSNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVRPV DNLL PSWAITLISVNGIFVICCL THCFAPRCRERRNERLRRESVHPV (241)(241)(241)(241)(241)241) 241) (241)(241)(241)241) (241) (241) (241)(241)(241)241) (241)241) (241)241) 241) 241) SEQ:087_R2_CTLA4BP-5x4-1f SEQ:088_R2_CTLA4BP-5x5-2e SEQ:089_R2_CTLA4BP-5x5-6e SEQ:090_R2_CTLA4BP-5x6-9d SEQ:091_R2_CTLA4BP-5x8-1f SEQ: 092_R2_CTLA4BP-5x9-12c SEQ:222_ctla5x9d10 SEQ:223_ctla5x6f6 SEQ:080_R2_CTLA4BP-5x2-7b SEQ:082_R2_CTLA4BP-5x3-10e SEQ:083_R2_CTLA4BP-5x3-11b SEQ:084_R2_CTLA4BP-5x3-6f SEQ:085_R2_CTLA4BP-5x4-11d SEQ:086_R2_CTLA4BP-5x4-12c SEQ:069_R1_CTLA4BP-5 SEQ: 072_R1_CTLA4BP-13 SEQ: 074_R2_CTLA4BP-5x2-10c SEQ:075_R2_CTLA4BP-5x2-11d SEQ:076_R2_CTLA4BP-5X2-12F SEQ:077_R2_CTLA4BP-5x2-2g SEQ:078_R2_CTLA4BP-5x2-3c SEQ:079_R2_CTLA4BP-5x2-4c SEQ:081_R2_CTLA4BP-5x2-8c SEQ:070_R1_CTLA4BP-7 SEQ:071_R1_CTLA4BP-11 SEQ: 073_R1_CTLA4BP-27

16/43

DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNGRLRRESVRPV

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DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVHPV

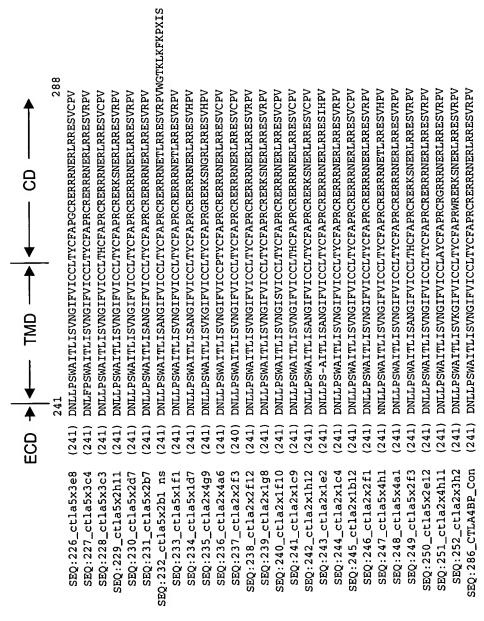


Fig. 3F

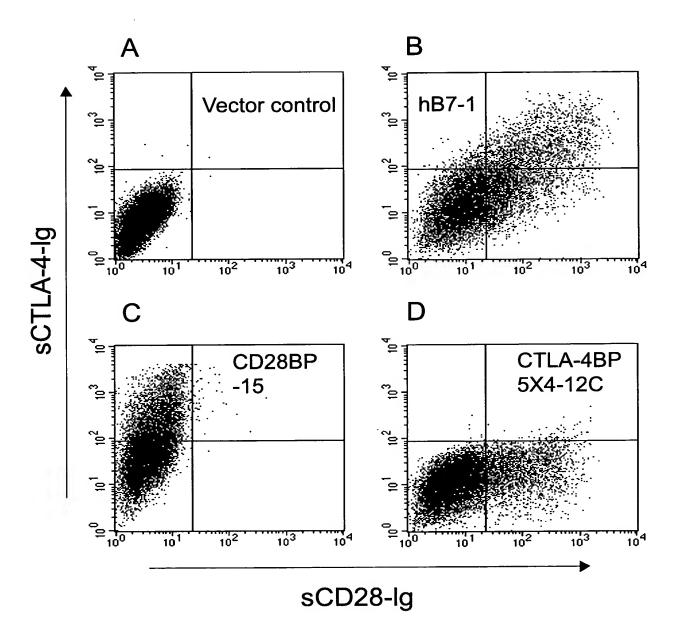


Fig. 4

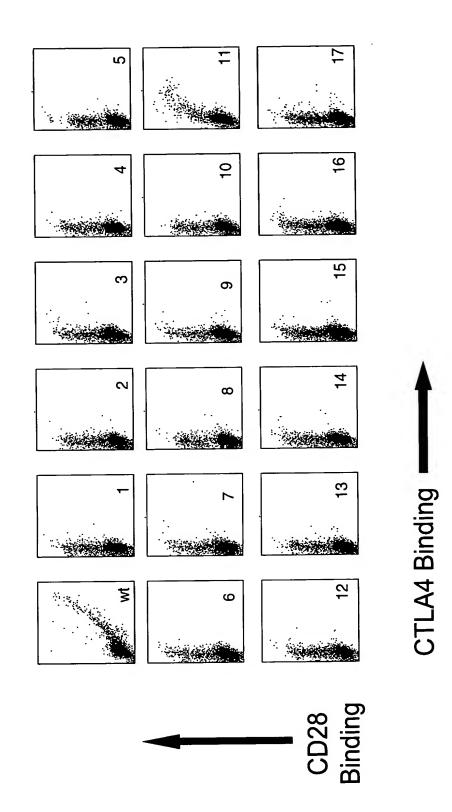
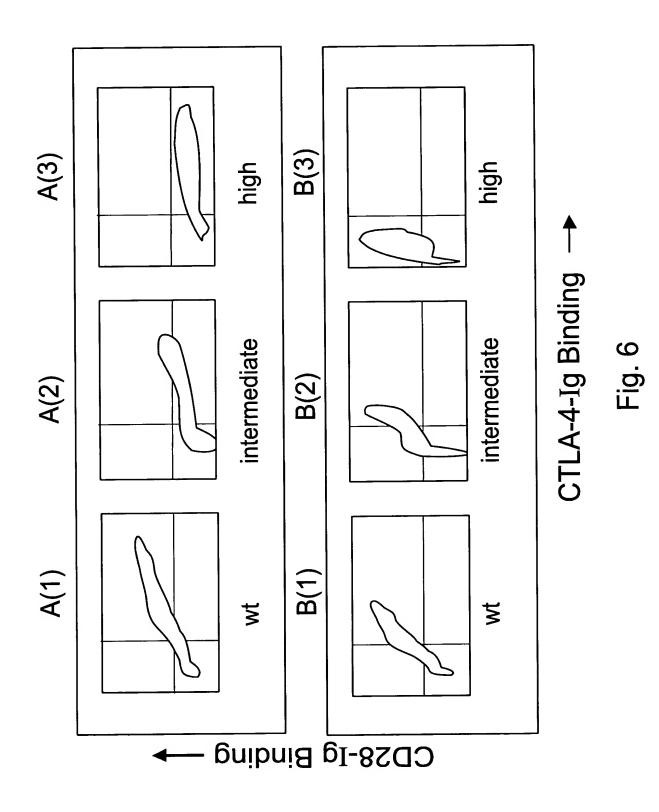


Fig. 5



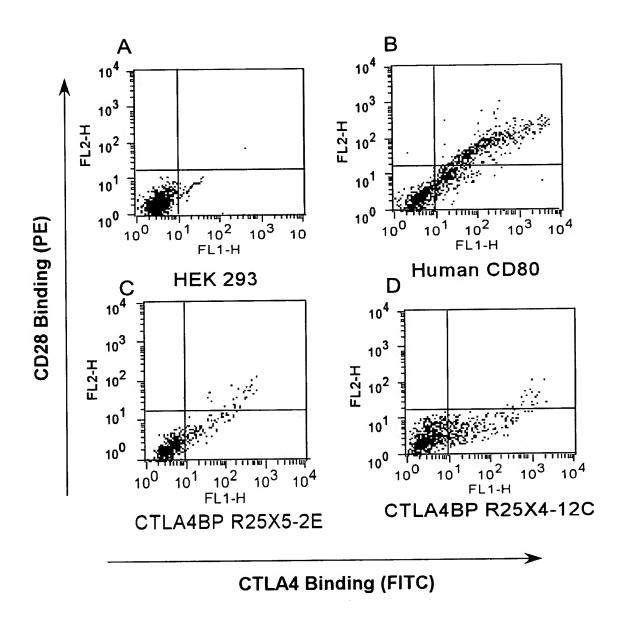


Fig. 7A-D

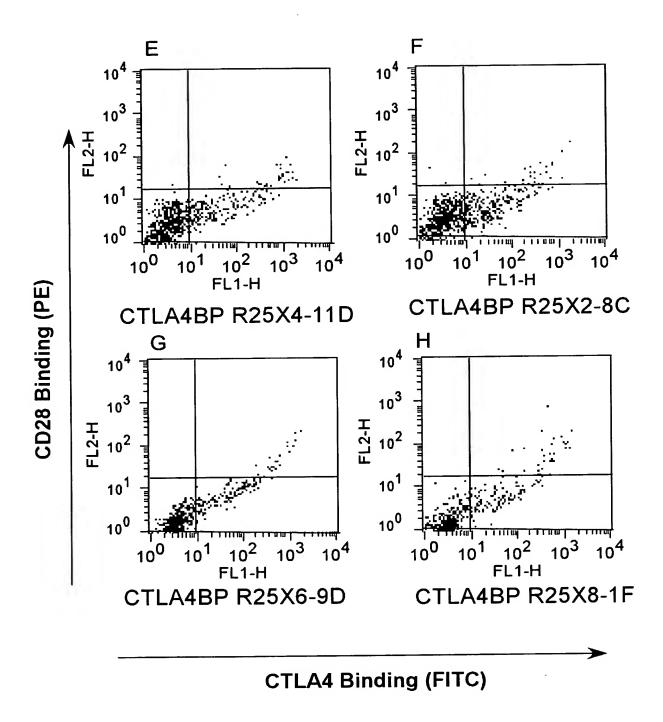


Fig. 7E-H

Fig. 8A

RIHWQKEKKMVLTMMSGDMNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAF KREHLAEVMLSVKADFPTPSISDFEIPPSNIRRIICSTSGGFPEPHLFWLENGEELNAINTTVSQ DPETELYTVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPKOEHFPDNLLPSWAITLISA MGHTRROGTSPSKCPYLŘFFQLLVLAGLSHFCSĠVIHVTKEVKEVATLSCGHNVSVEELAQT NGIFVICCLTYRFAPRCRERKŜNETLRRESVRPV CTLA-4BP

Fig. 8B

\$LRIYWQKD\$KMVLAILPGKVQVWPEYKNRTITDMNDNPRIVILALRPSD\$GTYTCVIQKPVLK ĞAŸKLEHLAŠVŘLMIRADFPVPŤINDLĞNPŠPNIRRLICSTSGGFPRPHLYWLENGEELNAŤNT MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEEL CD28BP

TVSQDPĞTELYMİSSELDFNVTNNHSİVCLIKYGELŞVŞQİFPWSKPKQEPPİDQLPFWVIIPVŞ

GALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG

rhesus/bab	%OS ☐	⊠ rabbi
M human	orangutan	rhesus

aboon papoon

000

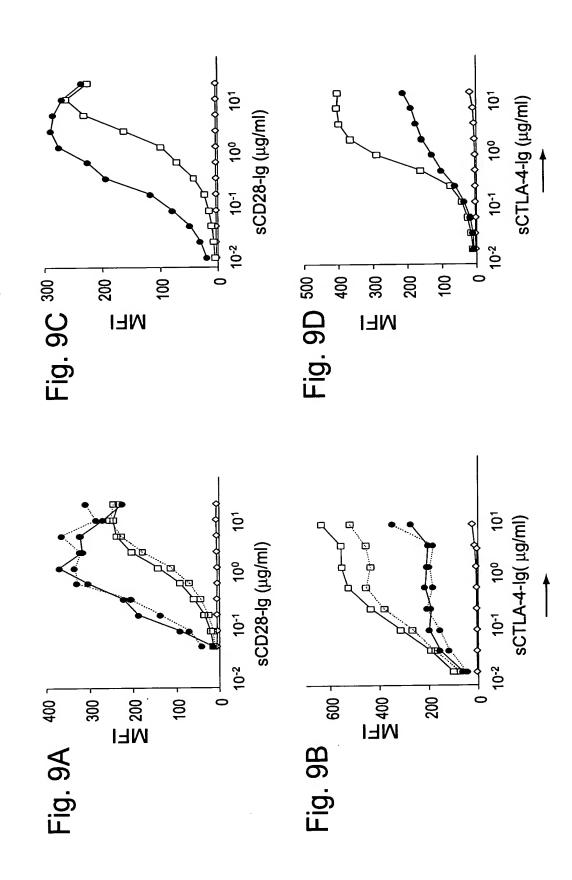
— CD28BP-Flag

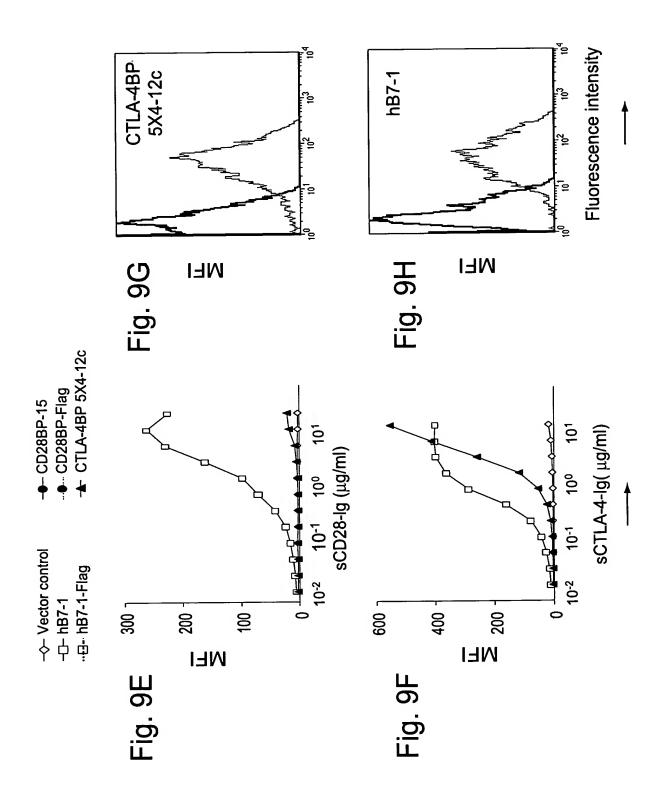
★ CTLA-4BP 5X4-12c

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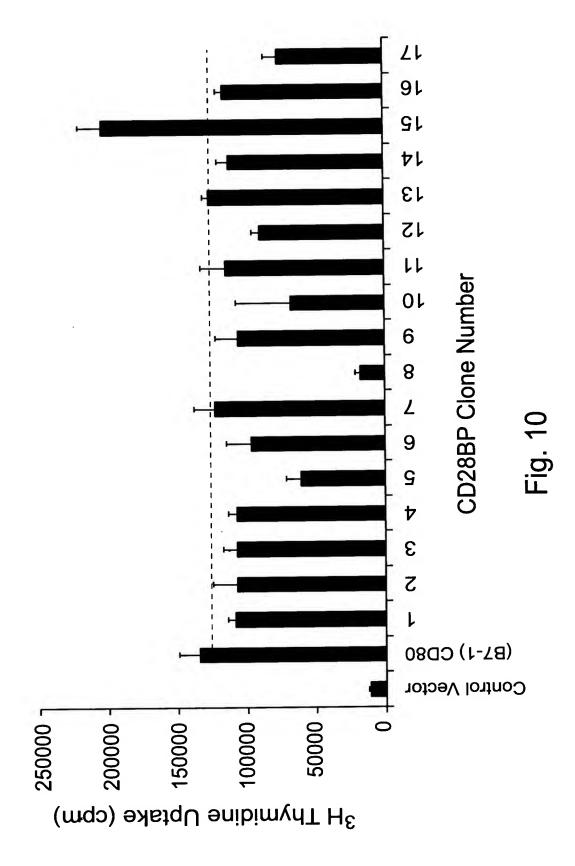
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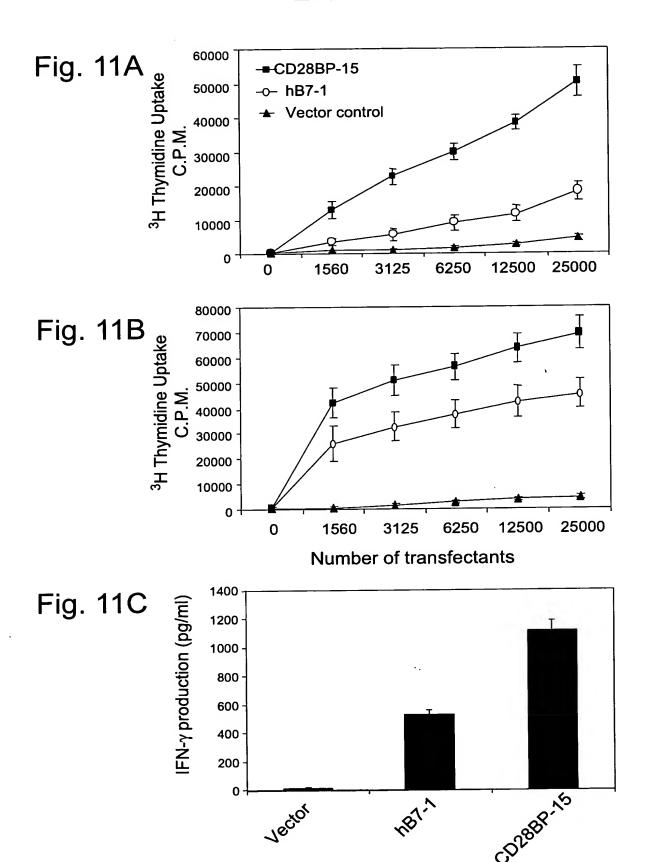
- CD28BP-15

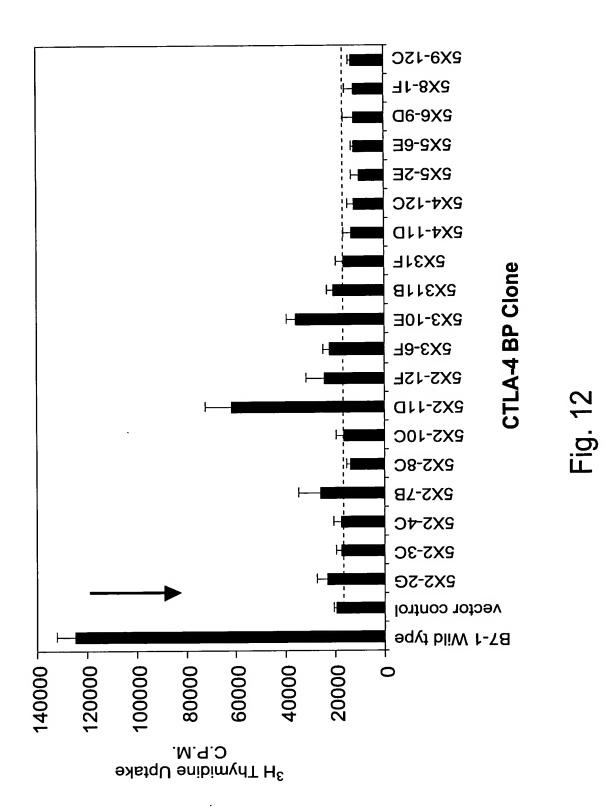


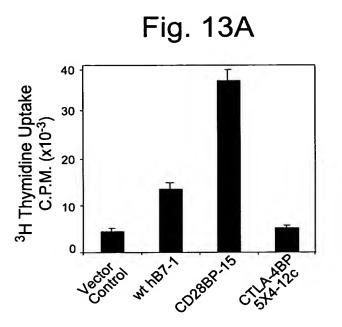


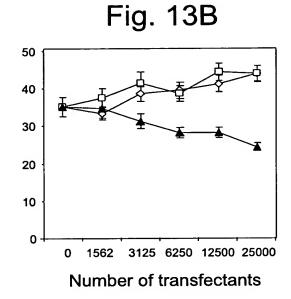


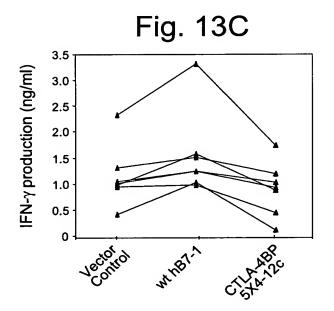












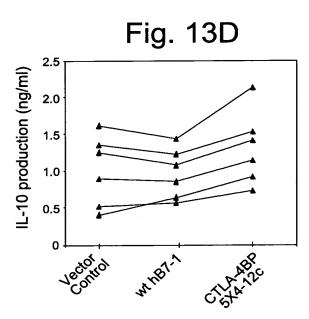


Fig. 14A

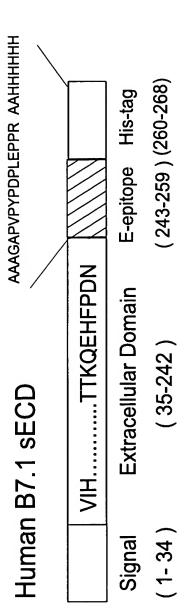
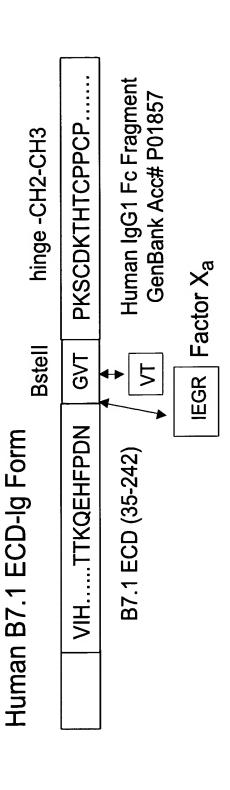


Fig. 14B



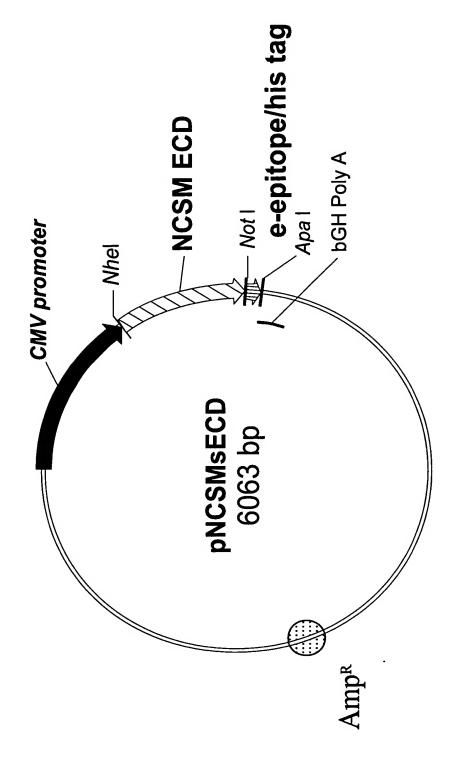


Fig. 15

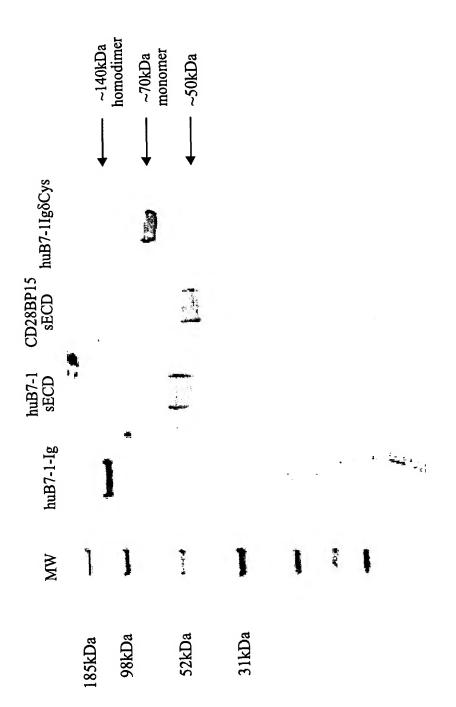


Fig. 16

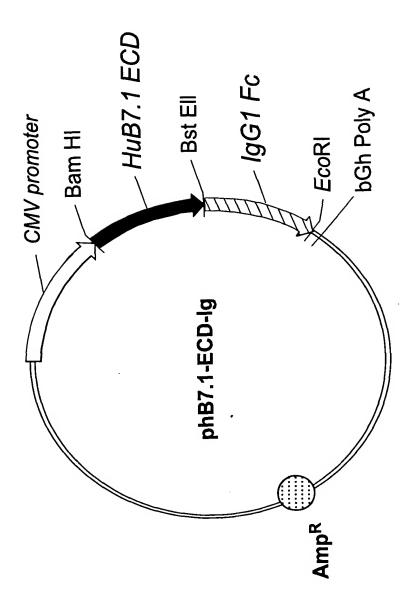


Fig. 17

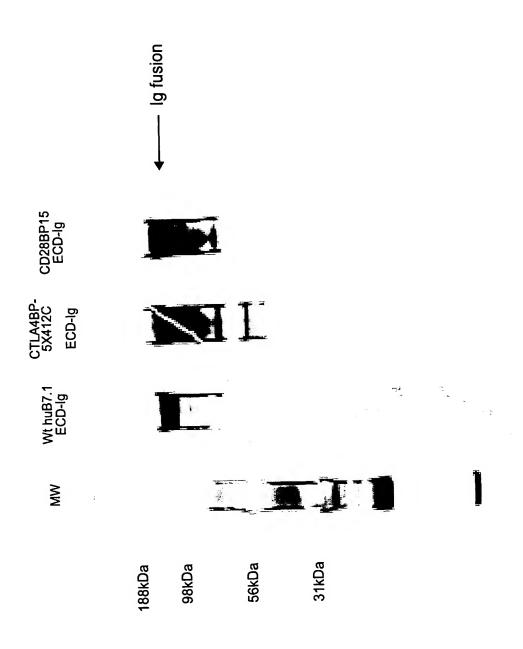


Fig. 18

Expression of CTLA-4BP-Ig and CD28BP-Ig Proteins

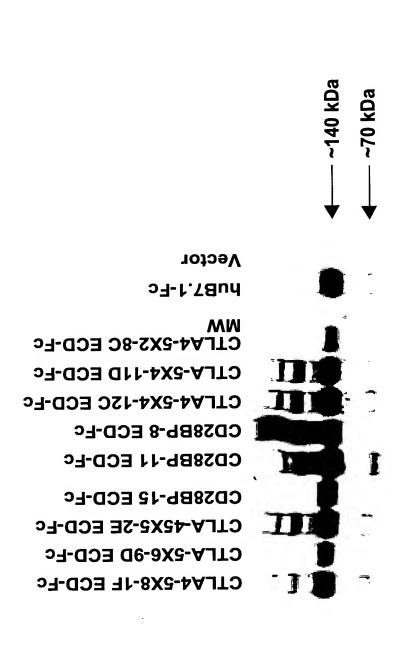


Fig. 20A

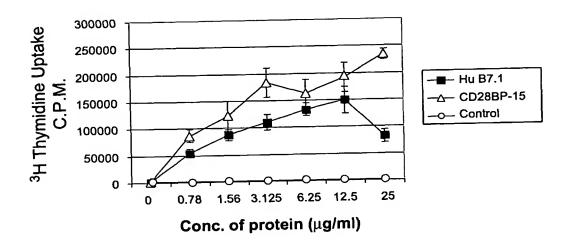


Fig. 20B

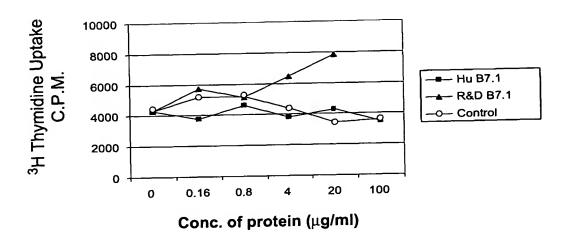
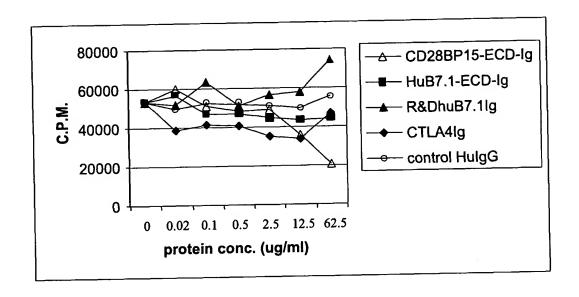
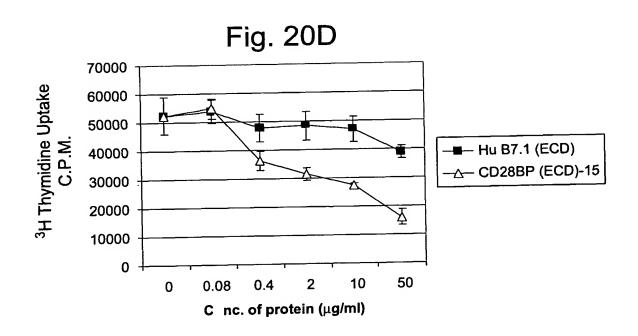


Fig. 20C





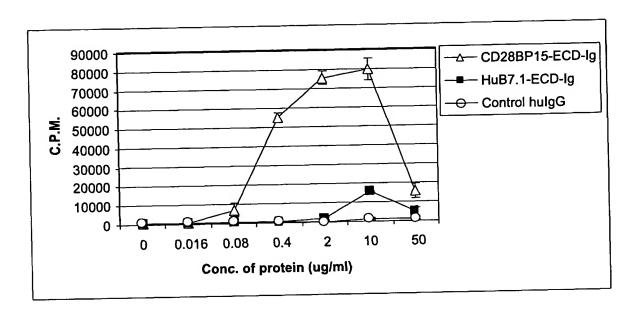


Fig. 20E

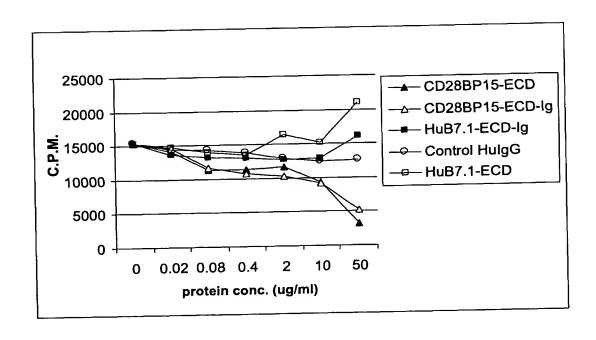


Fig. 20F

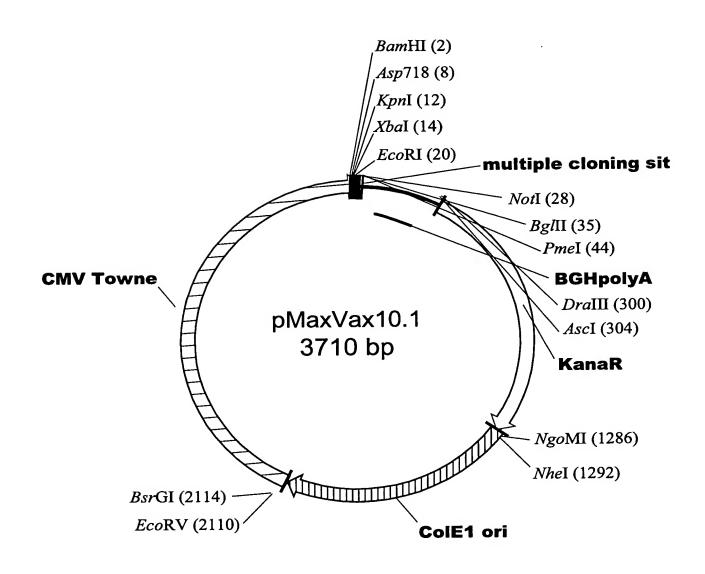


Fig. 21

Fig. 22A

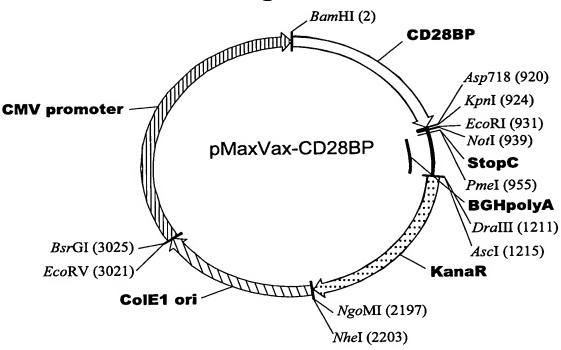
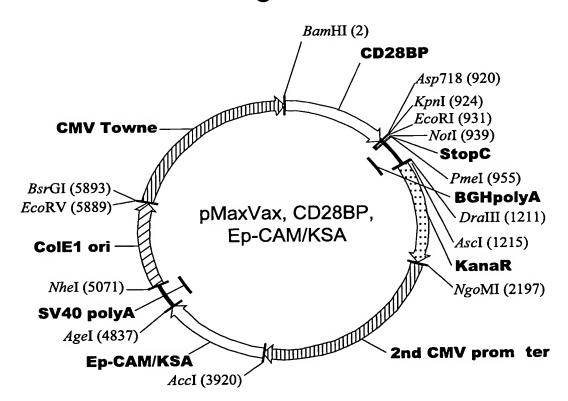


Fig. 22B



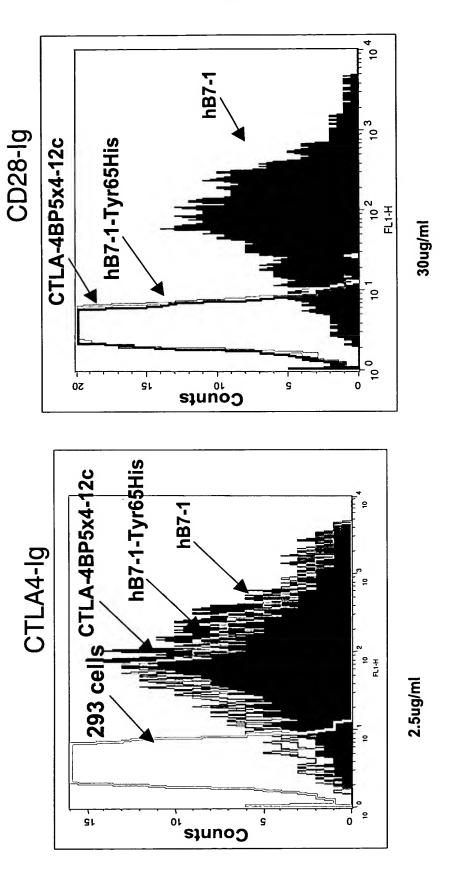


Fig. 23A

Fig. 23B

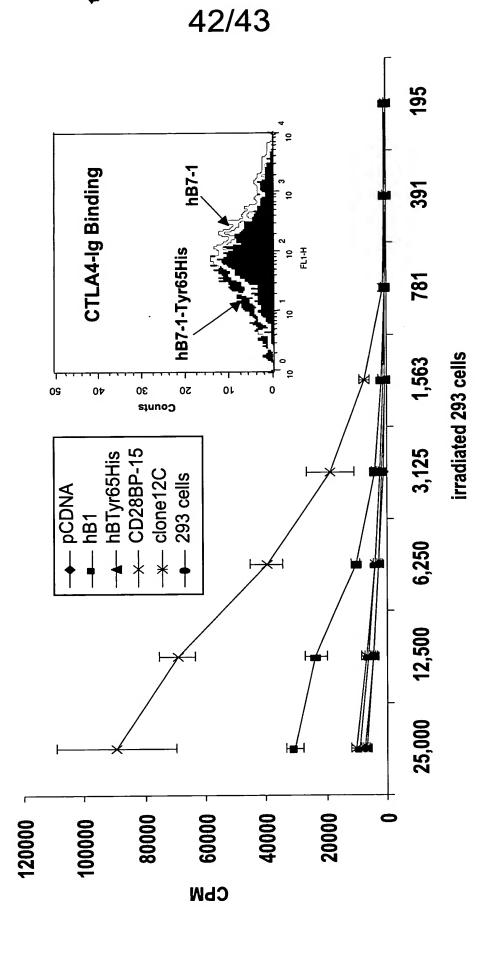


Fig. 24

Fig. 25A

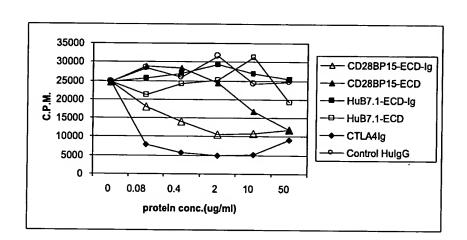


Fig. 25B

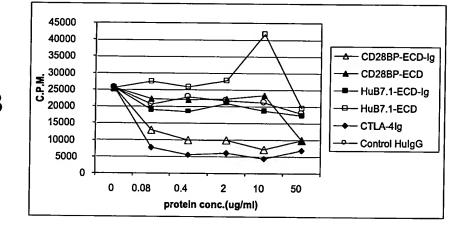
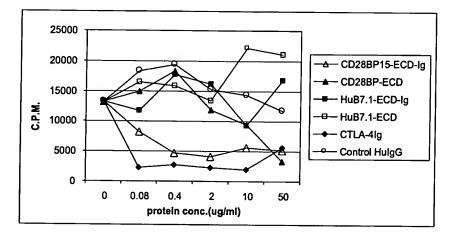


Fig. 25C



A